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Calcium-gated K^+ channels of the K_{Ca} 1.1- and K_{Ca} 3.1-type couple intracellular Ca^{2+} signals to membrane hyperpolarization in mesenchymal stromal cells from the human adipose tissue

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Abstract Electrogenesis in mesenchymal stromal cells (MSCs) remains poorly understood. Little is known about ion channels active in resting MSCs and activated upon MSC stimulation, particularly, by agonists mobilizing Ca²⁺ in the MSC cytoplasm. A variety of Ca2+-gated ion channels may couple Ca²⁺ signals to polarization of the plasma membrane. Here, we studied MSCs from the human adipose tissue and found that in cells responsive to ATP and adenosine with Ca²⁺ transients or exhibiting spontaneous Ca²⁺ oscillations, Ca²⁺ bursts were associated with hyperpolarization mediated by Ca²⁺-gated K⁺ channels. The expression analysis revealed transcripts for KCNMA1 and KCNN4 genes encoding for Ca²⁺-activated K⁺ channels of large (K_{Ca}1.1) and intermediate $(K_{Ca}3.1)$ conductance, respectively. Moreover, transcripts for the Ca²⁺-gated cation channel TRPM4 and anion channels Ano1, Ano2, and bestrophin-1, bestrophin-3, and bestrophin-4 were revealed. In all assayed MSCs, a rise in cytosolic Ca²⁺ stimulated K⁺ currents that were inhibited with iberiotoxin. This suggested that K_{Ca}1.1 channels are invariably expressed in MSCs. In ATP- and adenosine-responsive cells, iberiotoxin and TRAM-34 diminished electrical responses, implicating both K_{Ca}1.1 and K_{Ca}3.1 channels in coupling agonist-dependent Ca²⁺ signals to membrane voltage.

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Functional tests pointed at the existence of two separate MSC subpopulations exhibiting Ca²⁺-gated anion currents that were mediated by Ano2-like and bestrophin-like anion channels, respectively. Evidence for detectable activity of Ano1 and TRPM4 was not obtained. Thus, $K_{Ca}1.1$ channels are likely to represent the dominant type of Ca²⁺-activated K⁺ channels in MSCs, which can serve in concert with $K_{Ca}3.1$ channels as effectors downstream of G-protein-coupled receptor (GPCR)-mediated Ca²⁺ signaling.

Keywords Mesenchymal stromal cells $\cdot Ca^{2+}$ -activated K⁺ channels \cdot Patch clamp $\cdot Ca^{2+}$ imaging \cdot ATP \cdot Adenosine

Introduction

Mesenchymal stromal cells (MSCs) are defined as a heterogeneous cell population that includes adult multipotent cells capable of giving rise to differentiated cells of mesenchymal lines, such as osteoblasts, adipocytes, and chondrocytes [17, 30, 39]. First identified and isolated from the bone marrow, MSCs can now be obtained from a variety of other tissues, including the adipose tissue, umbilical cord blood, skin, and some others [6, 25, 33]. Although it is widely accepted that MSCs are crucial to tissue regeneration, molecular mechanisms controlling proliferation, migration, secretion, and differentiation of individual cells are still poorly understood.

Intracellular Ca^{2+} is a highly versatile second messenger involved in the regulation of almost all cellular functions [14, 49]. Particularly, spontaneous Ca^{2+} oscillations and Ca^{2+} transients evoked by local environmental cues can govern gene expression, cell proliferation and differentiation, and expansion of progenitor pools [2, 7, 31]. Thus, membrane receptors and signaling pathways associated with intracellular Ca^{2+} signaling may be critically important for the maintenance

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of a MSC population as well as for the regulation of MSC migration and differentiation.

Although the main function of ion channels is to control ion fluxes through the plasma membrane, they also can play a much less appreciated role in coordinating extracellular and intracellular signals associated with cell proliferation and differentiation [3, 5, 48, 57]. Ion channels can regulate exocytosis of autocrine and paracrine factors by setting Ca^{2+} influx directly or determining one indirectly via membrane potential [3, 7]. Ion channels are also capable of modulating cell proliferation and differentiation via flux-independent mechanisms, including association of channel proteins with membrane receptors [13, 66]. As a specific example, the Ca²⁺-activated cation channel TRPM4 is involved in differentiation of dental follicle stem cells, playing an inhibitory role in osteogenesis but being required for differentiation into adipocytes [42]. TRPM7 activity is required for bone marrow stem cell proliferation and viability [11]. The delayed rectifier K⁺ channels and Ca²⁺-activated K⁺ channels are required for MSC proliferation [16, 64]. Vascular smooth muscle cell proliferation critically depends on activity of large conductance Ca²⁺-gated K⁺ channel (K_{Ca}1.1 channels) [29].

As we found recently, the population of MSCs derived from the human adipose tissue is functionally heterogeneous in that it contains multiple subpopulation, each being specifically responsive to a particular agonist of G-protein-coupled receptors (GPCR), such as ATP, adenosine, noradrenalin, and some others [35]. These substances elicited Ca²⁺ transients in resting MSCs but weakly affected spontaneous Ca²⁺ oscillations observed in a small fraction of non-stimulated cells. Physiological significance of the agonist-dependent Ca²⁺ mobilization in MSCs and their oscillatory behavior remain largely unknown. Extracellular ATP, its downstream product adenosine, and related nucleotides are ubiquitous signaling molecules, operating in virtually all tissues and cells. In particular, MSCs are endowed with several purinergic receptors and ectonucleotidases and release extracellular purines presumably to mediate/promote growth/proliferation, pro- or anti-apoptotic processes, differentiation, and immunomodulation [52]. Here, we consider the immediate consequence of intracellular Ca²⁺ bursting, that is a Ca²⁺-dependent change in ion channel activity and provides evidence that in MSCs, Ca²⁺ signaling is associated with cell hyperpolarization that is largely mediated by Ca^{2+} -gated K⁺ channels of the K_{Ca}1.1 and K_{Ca}3.1 types.

Materials and methods

MSC isolation and culturing

Human MSCs were isolated from the subcutaneous adipose tissue of 15 healthy (i.e., without infectious, systemic diseases or malignancies) men from 32 to 60 years old. The body mass index (BMI) of individuals varied from 20 to 29. No clear correlation between BMI and physiological features of MSCs was found. All donors gave informed consent for harvesting their adipose tissue.

Isolation and characterization of cells were described earlier [35]. Briefly, the adipose tissue was extensively washed with two volumes of Hank's Balanced Salt Solution (HBSS) containing 5% antibiotic/antimycotic solution (HyClone), fragmented and then digested at 37 °C for 1 h in the presence of collagenase (200 U/ml, Sigma-Aldrich) and dispase (10 U/ ml, BD Biosciences). Enzymatic activity was neutralized by adding an equal volume of a culture medium (Advance Stem basal medium for human undifferentiated mesenchymal stem cells (HyClone)) containing 10% of Advance stem cell growth supplement (CGS) (HyClone) and 1% antibiotic/antimycotic solution and centrifuged at 200 g for 10 min. This led to the sedimentation of diverse cells, including MSCs, macrophages, lymphocytes, and erythrocytes, unlike adipocytes that remained floating. After removal of supernatant, a lysis solution (154 mM NH₄Cl, 10 mM KHCO₃, and 0.1 mM EDTA) was added to a cell pellet to lyse erythrocytes, and cell suspension was centrifuged at 200 g for 10 min. Sedimented cells were resuspended in the MSC culture medium and filtered through a 100 µm nylon cell strainer (BD Biosciences). As indicated by flow cytometry [35], after isolation and overnight pre-plating, the obtained cell population contained not only MSC cells that basically represented the most abundant subgroup but also admixed macrophages and lymphocytes. The two last cell subgroups were dramatically depleted by culturing for a week in the MSC culture medium and humidified atmosphere (5% CO₂) at 37 °C. The obtained MSC population was maintained at a subconfluent level (~80% confluency) and passaged using HyQTase (HyClone). For experiments, cells of the second to fourth passages were used.

Preparation of cells for patch clamping and Ca²⁺ imaging

Prior to physiological experiments, cells were maintained in a 12 socket plate for 12 h in the medium described above but without antibiotics. For isolation, cells cultured in a 1 ml socket were rinsed twice with the Versen solution (Sigma-Aldrich) that was then substituted for 0.2 ml HyQTase solution (HyClone) for 3-5 min. The enzymatic treatment was terminated by the addition of 0.8 ml culture medium to a socket. Next, cells were resuspended, and cell suspension was put into a tube for storage in the MSC culture medium at 4 °C for 6-8 h. When necessary, isolated cells were collected by a plastic pipette and plated onto a recording chamber. After sedimentation and attachment to the chamber bottom coated with the Cell-Tak adhesive (BD Biosciences), cells were loaded with Fluo-4 at room temperature (23-25 °C) by adding Fluo-4AM (4 µM) and Pluronic (0,02%) (all from Molecular Probes) to the bath solution (mM): 135 NaCl, 5 KCl, 1 MgCl₂, 2 CaCl₂, and 10

HEPES-NaOH (NaOH), pH 7.4. After the 20-min incubation, cells were rinsed several times with the bath solution.

Electrophysiology

Electrophysiological activity of MSCs was assayed with the patch clamp technique under the whole-cell (WC) mode or using the perforated patch approach. Ion currents were recorded, filtered, and analyzed using an Axopatch 200B amplifier, Digidata 1332A and MiniDigi 1A interfaces, and the pCLAMP 8 software (all from Molecular Devices). Cells were polarized either by serial voltage pulses of appropriate duration at a 10-20 mV step or by voltage ramp (1 mV/ms). The basic intracellular solution contained (mM) 140 KCl (CsCl), 1 MgCl₂, 0.1 EGTA, and 10 HEPES-KOH (KOH), pH 7.3. In certain cases, cells were dialyzed with the solution containing 440 nM free Ca²⁺ (mM): 140 CsCl, 1 MgCl₂, 7.9 CaCl₂ + 10 EGTA-CsOH, 10 HEPES-CsOH, and 2 ATP-Mg, pH 7.3. For WC or perforated patch recordings, 2 mM MgATP or amphotericin B (400 µg/ml) was added into the intracellular solution, respectively. The basic extracellular solution contained (mM) 135 NaCl, 5 KCl, 1 MgCl₂, 2 CaCl₂, and 10 HEPES-NaOH (NaOH), pH 7.4. When required, it was modified in that NaCl was replaced with Na-HEPES (low Cl^{-} solution). The recording chamber of nearly 75 µl and gravity-driven perfusion system have been described previously [34]. All chemicals were bath applied for 1 s and rinsed within 3 s. Buffers, salts, ATP, and adenosine were from Sigma-Aldrich; apamin, iberiotoxin, TRAM-34, CaCCinh-A01, and T16Ainh-A01 were from Tocris. Experiments were carried out at room temperature (22-24 °C).

Imaging

Cells loaded with Fluo-4 were imaged by using a fluorescent microscope Axioscope 2 equipped with an objective Plan-Neofluar 20×/0.50 (Zeiss) and ECCD camera LucaR (Andor Technology). Apart from a transparent light illuminator, the microscope was equipped with a handmade system for epiillumination via an objective. Fluo-4 fluorescence was excited by using a computer-controllable light emitting diode (LED) LZ1-00B700H (Ledengin). LED emission was filtered with an optical filter ET480/20× (Chroma Technology). Fluo-4 emission was collected at 535 ± 25 nM by using emission filter ET535/50m (Chroma Technology). Serial fluorescent images were captured every second and analyzed with the Imaging Workbench 6 software (INDEC). The deviation of cytosolic Ca²⁺ from the resting level in an individual cell was quantified by the relative change in Fluo-4 fluorescence $\Delta F/$ F_0 , where $\Delta F = F - F_0$; F is the instant intensity of cell fluorescence and F_0 is the intensity of cell fluorescence obtained in the very beginning of a recording and averaged over a 20-s interval. In certain cases, cells preloaded with Fluo-4 were patch clamped, and their electrophysiological characteristics were analyzed simultaneously with intracellular Ca²⁺ monitoring.

RT-PCR

Total RNA was extracted from a sample containing $10^5 - 10^6$ MSCs by using the RNeasy Mini Kit (Qiagen). Isolated RNA was treated with DNase I (Ambion) and reverse transcribed with PrimeScript reverse transcriptase (TaKaRa) and random hexamer primers, following the manufacturer's instruction. Obtained cDNA served as a template for PCR with genespecific primers that were designed to recognize sequences of all known splice variants of human genes encoding Ca²⁺gated K⁺ channels, bestrophins, and anoctanins as well as three marker genes for MSCs, including CD73, CD90, and CD105 (Supplementary Materials, Table S1). The expected sizes of PCR products were as follows (in bp): KCNMA1-446. KCNN1-316. KCNN2-445. KCNN3-268. KCNN4-330; CD73-266, CD90-344, CD105-317; TRPM4-367, TRPM5-273; Ano1-295, Ano2-329, Ano3-444, Ano4-311, Ano5-252, Ano6-262, Ano7-321, Ano8-364, Ano9-367, Ano10 and Ano332; Best1-282, Best2-300, Best3-198, Best4-316. The PCR products were verified by direct sequencing (Evrogen, Moscow).

Results

In order to probe functional coupling of Ca^{2+} signals to membrane polarization, a number of MSCs were concurrently assayed with the Ca^{2+} imaging and the patch clamp technique. In designated experiments, we loaded MSCs with Fluo-4 and searched for cells that exhibited spontaneous Ca²⁺ oscillations or were responsive to ATP or adenosine, the purinergic agonists shown to mobilize intracellular Ca^{2+} in MSCs [19, 35, 67]. If generated spontaneous or agonist-induced Ca²⁺ signals, a cell was then examined with the patch clamp technique. It should be noted that patch clamping frequently stopped Ca²⁺ oscillations in wavering cells and also rendered purinergic MSCs irresponsive to ATP and adenosine for a currently undetermined reason. Perhaps, being unavoidable for gigaseal formation, the mechanical disturbance of an assayed cell destroyed intracellular machinery responsible for Ca²⁺ signaling. Because of this interfering phenomenon, most of oscillating and purinergic MSCs could not be effectively assayed with the patch clamp technique and Ca²⁺ imaging in concurrent recordings. Eventually, we succeeded in conclusive recordings from seven MSCs with oscillating cytosolic Ca²⁺ and found that in all of them, membrane voltage varied synchronously with cytosolic Ca^{2+} (Fig. 1a). Given that a rise in cytosolic Ca²⁺ was accompanied by cell hyperpolarization, Ca²⁺-dependent stimulation of K⁺ channels or inhibition of



Fig. 1 Oscillations of intracellular Ca^{2+} are associated with a periodic increase in K⁺ conductance. **a** Representative example (n = 7) of concurrent monitoring of intracellular Ca^{2+} (*upper panel*) and membrane voltage (*bottom panel*) in MSC loaded with Fluo-4. *Upper panel*, the data are presented as $\Delta F/F_0$, where $\Delta F = F - F_0$; *F* is the instant intensity of cell fluorescence, and F_0 is the intensity of cell fluorescence obtained in the very beginning of the recording and averaged over a 20-s interval. *Bottom panel*, membrane voltage synchronously recorded under zero-current clamp. The recording was interrupted at the moments *I* and *2* to generate I-V curves under the voltage clamp mode. In this case, an

cationic and/or Cl⁻ channels was obligatory for this phenomenon. To probe into ionic specificity of responsible channels, we generated current-voltage (I-V) curves when Ca²⁺ reached the lowest and maximal levels. Since input resistance varied from cell to cell within an order of magnitude, current injection produced an unpredictable voltage response in a given cell assayed under the current clamp mode. We therefore used the following protocol. Basically executed under zero-current clamp, a recording was switched to voltage clamp for 3-5 s to generate an I-V curve with a voltage ramp (1 mV/ms) polarizing a cell from -100 to 80 mV, and then, the acquisition was returned to the current clamp (Fig. 1a, bottom panel, triangles 1 and 2). The representative I-V curves shown in Fig. 1b (left panel) clearly indicated that a Ca²⁺ transient in the MSC cytoplasm stimulated a marked increase in membrane conductance and shifted a reversal potential of an integral current to the left by 30-40 mV. The Ca²⁺-dependent current, which was determined as a difference between integral currents measured at minimal and maximal Ca²⁺ (Fig. 1b, right panel), reversed at nearly -78 ± 1.3 mV on average (n = 9). This value is quite close to the Nernst potential for K⁺ ions that is equal to -85 mV at the [140 mM]_{in}/[5 mM]_{out} gradient and 22 °C. It thus appeared that the periodic hyperpolarization observed in MSCs with spontaneous Ca²⁺ oscillations (Fig. 1a) was

assayed cell was held at -40 mV and polarized by a voltage ramp (1 mV/ ms) between -100 and 80 mV. **b** *Left panel*, I-V curves generated at the moments *I* and *2* as indicated in the *bottom panel* in (**a**). *Right panel*, the I-V curve of Ca²⁺-dependent current determined as a difference of I-V curves 2 and 1 presented in the *left panel*. The current reversed at nearly -80 mV (*arrow*), suggesting Ca²⁺-activated K⁺ channels to be responsible. The perforated patch approach was employed. The patch pipette was filled with 140 mM KCl + 400 µg/ml amphotericin B; the bath solution contains 135 mM NaCl +5 mM KCl

mainly associated with stimulation of Ca^{2+} -gated K⁺ channels by Ca^{2+} bursts.

Expression of Ca²⁺-gated ion channels in MSCs

The family of Ca²⁺-activated K⁺ channels is functionally subdivided into small (K_{Ca}2.x), intermediate (K_{Ca}3.1), and large (K_{Ca}1.1) conductance channels [4, 56, 61]. In mammalians, a single gene KCNMA1 encodes the pore-forming α subunit of K_{Ca}1.1 channels [50], three genes, including KCNN1, KCNN2, and KCNN3 encode K_{Ca}2.x channels, while KCNN4 encodes the K_{Ca}3.1 channel [56, 65]. By using conventional RT-PCR and gene-specific primers, we analyzed expression of the abovementioned genes in MSCs. It turned out that all RNA preparations (n = 5) isolated from MSC colonies of nearly 10⁶ cells contained transcripts for KCNMA1 and KCNN4, while KCNN1, KCNN2, and KCNN3 transcripts were undetectable (Fig. 2a). These findings indicated that Ca²⁺-activated K⁺ currents in MSCs (Fig. 1) were presumably mediated by K_{Ca} 1.1 and/or K_{Ca} 3.1 channels. Apart from specialized K⁺ channels mentioned previously, Ca²⁺ ions also gate the cation channels TRPM4 and TRPM5 [21] as well as anion channels formed by certain channel subunits from the anoctamin (TMEM16) and



Fig. 2 RT-PCR analysis of expression of Ca^{2+} -gated channels in MSCs. **a** Detected amplicons of expected sizes (bp) correspond to transcripts for the *KCNMA1* (446) and *KNCK4* (330) genes encoding for K_{Ca}1.1 and K_{Ca}3.1 channels, respectively, as well as for the marker genes *CD73* (266), *CD90* (344), and *CD105* (317). Products of expected sizes for *KCNN1* (316), *KCNN2* (445), and *KCNN3* (268) were not detectable. **b** Detection of transcripts for anoctamins. Fragments of expected sizes were obtained for *Ano1* (295), *Ano2* (329), *Ano3* (444), *Ano4* (311), *Ano5* (252), *Ano6* (262), *Ano8* (364), and *Ano10* (332). Transcripts for *Ano7*

bestrophin families. Among them, Ano1 (TMEM16A), Ano2 (TMEM16B), and Best1 have been reported to form Ca²⁺-activated anion channels in a variety of cells [37, 53]. These Ca²⁺-dependent cation and anion channels could also couple intracellular Ca²⁺ signals to plasma membrane polarization. We therefore analyzed their expression in MSCs and detected transcripts for Ano1 and Ano2 and several other anoctamins as well as for bestrophin-1, bestrophin-3, and bestrophin-4 (Fig. 2c, d).

Ion currents regulated by intracellular Ca²⁺ in MSCs

Next, we tried to reveal functional activity of the Ca²⁺-gated ionic channels that were identified in MSCs at the messenger RNA (mRNA) transcript level (Fig. 2). Note that intracellular Ca^{2+} oscillated spontaneously (Fig. 1a) in a small (3–5%) MSC subpopulation so that the vast majority of assayed cells exhibited relatively stable resting potentials ranging between -10and -40 mV with 140 mM KCl in the pipette and 135 mM NaCl + 5 mM KCl in the bath. When treated with Ca^{2+} ionophore ionomycin (5 µM), all MSCs assayed in this series (n = 51) were hyperpolarized by 20–40 mV. Iberiotoxin (IbTX) (100 nM), a specific blocker of K_{Ca}1.1 channels, reversed the ionomycin effect partly or completely (Fig. 3a). Consistent with activity of K_{Ca}1.1 channels, large IbTX-sensitive, outwardly rectifying, voltage-gated currents were seen in cells, wherein ionomycin sufficiently elevated cytosolic Ca²⁺ (Fig. 3b, c). In contrast, ionomycin effects were not reversed by the specific $K_{Ca}3.1$ channel blocker TRAM-34 (1 μ M) in most (88%) assayed MSCs (Fig. 3d, e), suggesting negligible

(321) and Ano9 (367) were undetectable. **c** Detection of transcripts for bestrophins and TRPM channels. Fragments of expected sizes were obtained for Best1 (282), Best4 (316), and TRPM4 (367). In the case of Best3, transcript variant 2 (NM 152439.3) (198) was found, while transcript variants 1, 3, and 4 (351) as well as Best2 (300) and TRPM5 (273) were not detected. The molecular weight markers (M) were GeneRuler 100 bp DNA Ladder (Fermentas). The agarose gels (1.3%) were stained with ethidium bromide. No specific signals were detected in the no RT control

contribution of $K_{Ca}3.1$ channels to the ionomycin-induced hyperpolarization of these cells. Nevertheless, among 51 assayed MSCs hyperpolarized by 5 µM ionomycin, six cells (12%) were significantly or almost completely repolarized by 1 µM TRAM-34, while 100 nM IbTX elicited only small additional depolarization (Fig. 3f). In all these cases, TRAM-34 exerted a small decrease in membrane conductance that was detectable at negative voltages but masked at positive potentials by large and nosy outward currents mediated by $K_{Ca}1.1$ channels (Fig. 3g). Together, the abovementioned findings led us to the conclusion that $K_{Ca}1.1$ channels represent the main type of Ca^{2+} -gated K⁺ channels that were functional in the vast majority, if not each of MSCs, while ionomycin-induced activity of $K_{Ca}3.1$ channels was characteristic of a small MSC subpopulation.

Because large and noisy currents mediated by K_{Ca}1.1 channels greatly complicated reliable identification of Ca²⁺-dependent anionic and cationic channels, we dialyzed a number of hAD-MSCs with a CsCl-based intracellular solution to completely suppress Ca²⁺-activated K⁺ currents. When perforated patch approach was employed, 5 µM ionomycin elicited marked inward currents in 12 out of 45 cells (27%) held at -60 mV and increased their conductance (Supplementary Materials, Fig. S1). In all cases, ionomycin-induced currents in Cs⁺-dialyzed cells were suppressed by the common inhibitor of Ca²⁺-gated anionic channels CaCCinh-A01 (CaCCinh) (100 µM) (Fig. S1) and several blockers of anionic channels such as SITS (500 μ M) and 9-AC (2 mM) (not shown). In contrast, assayed cells were negligibly sensitive to 9phenanthrol (300 nM) (Fig. S1), a specific TRPM4 blocker [22]. Thus, with CsCl in the pipette and NaCl in the bath,



Fig. 3 Ionomycin hyperpolarizes MSC by stimulating Ca^{2+} -gated K⁺ channels. **a** Representative (n = 51) recording of membrane voltage in MSC stimulated by 5 μ M ionomycin and treated with 100 nM iberiotoxin (IbTX) as indicated by the *horizontal lines above the experimental trace*. **b** Families of integral currents elicited by voltage pulses in the same cell in control, with 5 μ M ionomycin in the bath (*middle panel*), and in the presence of 100 nM iberiotoxin (IbTX) as indicated the moments when current values were measured to generate I-V curves shown in (**c**). The assayed cell was held at -60 mV and polarized by 140-ms voltage pulses between -100 and 80 mV. **d** Representative zero-current clamp recording (n = 45) from

ionomycin stimulated anionic currents, while no evidence for TRPM4 activity in MSCs was obtained. Perhaps, TRPM4-positive cells represent a minor group in a MSC population.

Reportedly, anion currents mediated by Ano1 and Ano2 channels activate slowly and exhibit marked outward rectification at a physiologically moderate level of intracellular Ca²⁺, while at micromolar Ca²⁺, those become almost ohmic [12,

MSC negligibly sensitive to TRAM-34. A cell was stimulated by 5 μ M ionomycin and treated with 1 mM TRAM-34 and 100 nM IbTX as indicated. The recording was interrupted at the moments 1-4 to generate I-V curves by voltage ramp. **e** I-V curves generated in (**d**) at the moments 1-4 indicate that in a given cell, ionomycin stimulated an outwardly rectifying K⁺ current that was blocked by IbTX, pointing at K_{Ca}1.1 channels to be responsible. **f** Representative recording (n = 6) from a cell sensitive to both TRAM-34 and IbTX. **g** I-V curves generated at the moments 1-4 in (**f**) indicate that both K_{Ca}1.1 and K_{Ca}3.1 channels were responsible for hyperpolarization of a given cell by ionomycin. The recording conditions were as in Fig. 1

46]. Thus, at high intracellular Ca²⁺, Ano1, and Ano2 channels are poorly distinguishable at the level of integral currents from bestrophin channels, which mediate almost instantly activating and weakly voltage-dependent currents [23, 41]. Yet, at the moment, no subtype-specific blockers of anoctamin and bestrophin channels are available, excluding T16inh-A01 (T16), an inhibitor of Ano1 channels [15]. Because the accurate control of intracellular Ca^{2+} with ionomycin is hardly possible, we dialyzed MSCs via the patch pipette containing Ca^{2+} -EGTA buffer to maintain 440 nM free Ca^{2+} in CsCl-based intracellular solution. At this concentration of intracellular Ca^{2+} , Ano channels were expected to remain slowly activating, while bestrophin channels should have endowed the plasma membrane with an ohmic conductance.

Overall, we succeeded in sufficiently stable whole-cell (WC) recordings from 48 MSCs held at -60 mV, and in 35 cells (73%), the introduction of 440 nM Ca²⁺ affected insignificantly their resting currents and I-V characteristics that remained nearly linear during dialysis (not shown). This suggested that neither Ca²⁺-gated anionic channels nor TRPM4 were detectably active in such cells under our recording conditions. Meanwhile, the dialysis greatly increased resting currents in 13 cells (27%) and produced dramatic transformation of their I-V curves (Figs. 4 and 5). Among them, 5 cells reacted to the introduction of 440 nM Ca²⁺ with gradually increasing resting current (Fig. 4a) that was associated with the appearance of slowly activating voltage-gated (VG)

currents with strong outward rectification and reversal potential closed to zero voltage (Fig. 4b, c). These Ca²⁺-dependent currents were significantly suppressed by 100 μ M CaCCinh (Fig. 4a, b). Our findings indicated that predominantly, anionic channels were responsible for the slowly activating currents stimulated by intracellular Ca²⁺ at the CsCl/NaCl gradient. Note that these Ca²⁺-gated anionic currents were insensitive to 10 μ M T16 (Fig. 4a, b).

Unexpectedly, we faced the interfering artifact effect of CaCCinh on ion permeability of MSCs: in 23 out of 32 cells treated with CaCCinh, the compound stimulated VG currents at concentrations that were usually sufficient to inhibit Ca²⁺-gated anion currents (Supplementary Materials, Fig. S2). With 140 mM CsCl in the recording pipette and 140 mM NaCl in the bath, CaCCinh-activated currents reversed between -90 and -80 mV pointing at Cs⁺ ions as the main charge carriers. Given this finding and that CaCCinh-activated currents disappeared when 20 mM tetraethylammonium (TEA) was added to the bath (n = 14) (Fig. S2), we inferred that CaCCinh directly or indirectly stimulated VG K⁺ channels well permeable



Fig. 4 Slowly activating Ca²⁺-gated anion currents. **a** Evolution of a *WC* current in a cell held at -60 mV during introduction of 440 nM Ca²⁺. The Ca²⁺-dependent current was insensitive to the Ano1 blocker T16 (10 μ M) but suppressed by CaCCinh (100 μ M), an inhibitor of Ca²⁺-gated anion channels. At the moments *l*-5, the cell was polarized by 250-ms voltage pulses from -100 to 90 mV to generate I-V curves; the associated current transients were removed from the current trace. The disruption of a cell-attached patch was performed at the moment marked as *WC*. The patch pipette contained 140 mM CsCl and 7.9 mM CaCl₂ + 10 mM EGTA

(440 nM free Ca²⁺); 140 mM NaCl was present in the bath. **b** Families of integral currents elicited by voltage pulses and recorded at the corresponding moments 1-5 shown in (**a**). **c** Left and middle panels, I-V curves of the integral currents 1-5 shown in (**b**) and measured at the moments indicated by the symbols above the current traces. Right panel, the I-V curve of the Ca²⁺-dependent slowly activating current inhibitable with CaCCinh that was calculated as a difference between the WC currents recorded at the moments 3 and 5

Because TEA inhibited the stimulatory effects of CaCCinh completely (Fig. S2), we calculated a Ca²⁺-gated anion current as a difference between a WC current recorded when a resting current reached the plateau (Fig. 4a, moment 3) and one measured in the presence of 100 μ M CaCCinh +20 mM TEA (Fig. 4a, moment 5). By this approach, CaCCinh-sensitive Ca²⁺-dependent currents were characterized by I-V curves with strong outward rectification (Fig. 4c, right panel). Kinetically and by the voltage dependence, the responsible Ca²⁺-gated anionic channels were similar to Ano1/Ano2 channels rather than to bestrophin channels. The insignificant effect of T16 (Fig. 5) argues against a contribution of Ano1, thus implicating Ano2 channels in mediating slowly activating Ca²⁺-gated anion currents observed in MSCs.

Apart from slowly activating currents (Fig. 4), the introduction of 440 nM Ca²⁺ stimulated activating currents in 13 MSCs almost instantly (Fig. 5). These Ca²⁺-dependent currents were suppressed by 100 μ M CaCCinh (Fig. 5a, b) as well as by 500 μ M SITS and 2 mM 9-AC (not shown), suggesting anionic selectivity of responsible Ca²⁺-gated channels. For the reason mentioned previously, a Ca²⁺-dependent anion current was evaluated in the presence of 20 mM TEA and determined as a difference between WC currents recorded in control (Fig. 5a, moment 4) and after the addition of 100 μ M CaCCinh (Fig. 5a, moment 5). As exemplified by Fig. 5c (right panel), CaCCinh-suppressible currents reversed at nearly zero voltage and exhibited weak outward rectification. The immediate activation kinetics (Fig. 5b) and subtle voltage dependence of these Ca²⁺-gated anionic currents (Fig. 5b, c, right panel) point at bestrophin channels to be largely responsible for anion flux.

Involvement of K_{Ca} 1.1 and K_{Ca} 1.1 channels in transduction of ATP and adenosine

Although the ionomycin treatment of MSCs rendered Ca^{2+} dependent K⁺ and anionic channels active (Figs. 3, 4, and 5), underlying Ca^{2+} signals might be much higher, more lasting, and/or inappropriately localized compared to Ca^{2+} transients elicited by natural agonists. In certain



Fig. 5 Instantly activating Ca^{2+} -gated anion currents. **a** Introduction of 440 nM Ca^{2+} stimulated inward current in a cell held at -60 mV and resulted in an increase in membrane conductance as shown in (**b**). At the moments *l*-5, the cell was polarized by 250-ms voltage pulses from -100 to 90 mV to generate I-V curves; the associated current transients were removed from the current trace. **b** The recording conditions were as Fig. 4. **b** Families of integral currents elicited by voltage pulses and

recorded at the corresponding moments l-5 shown in (a). c Left and middle panels, I-V curves of the integral currents 1–5 shown in (b) and measured at the moments indicated by the symbols above the current traces. Right panel, the I-V curve of the Ca²⁺-dependent, CaCCinh-suppressible, instantly activating current that was calculated as a difference between WC currents recorded at the moments 4 and 5

experiments, we therefore checked whether Ca²⁺-gated channels could be stimulated by the purinergic agonists ATP and adenosine, both being previously documented by us to mobilize Ca²⁺ in MSCs [35]. To simultaneously monitor intracellular Ca²⁺ and electrophysiological responses to the agonists, MSCs were loaded with Fluo-4. As with oscillating MSCs (Fig. 1), assayed cells were stimulated with 10 µM ATP or 1 µM adenosine; agonist-sensitive MSCs were identified by Ca²⁺ responses, and then those were patched. Overall, we succeeded in conclusive Ca²⁺ imaging/electrophysiological recordings from nine MSCs that remained ATP responsive after patching. These cells were usually clamped at zero current, and recordings of membrane voltage were briefly switched to voltage clamp for generating I-V curves when necessary. In all these cells, a short application of 10 μ M ATP elicited marked Ca²⁺ bursts (Fig. 6a, upper panel) that were accompanied by transient membrane hyperpolarization (Fig. 6a, bottom panel, c). By generating I-V curves prior and during ATP application when intracellular Ca²⁺ peaked (Fig. 6b, left panel), it was demonstrated that electrophysiological responses to ATP largely resulted from an increase in outwardly rectifying K⁺ conductance (Fig. 6b, right panel). In the presence of 1 µM TRAM-34, voltage responses to 10 µM ATP were nearly twice lower compared to control, although the nucleotide elicited similar Ca²⁺ transients in both cases (Fig. 6a, e). When both TRAM-34 (1 μ M) and IbTX (100 nM) were added to the bath, hyperpolarizing ATP responses were completely but reversibly inhibited (Fig. 6a, c). Note that 10 µM ATP insignificantly affected a membrane conductance in the presence of the blockers of K_{Ca}3.1 and K_{Ca}1.1 channels (Fig. 6d, middle panel). This observation pointed at a negligible contribution of P2X receptors to electrophysiological responses of MSCs to ATP.

Next, we focused on adenosine-sensitive MSCs and obtained overall eight cells with retained capability to respond to adenosine with Ca²⁺ transients after gigaseal formation. In all cases, briefly applied adenosine $(1 \mu M)$ triggered not only Ca²⁺ bursts but also marked hyperpolarizing voltage responses (Fig. 7a). As for ATP responses (Fig. 6a, b), adenosine stimulated outwardly rectifying currents that reversed at nearly -80 mV (Fig. 7b), the observation implicating Ca²⁺gated K⁺ channels in mediating adenosine-dependent MSC hyperpolarization. While both Ca²⁺ and voltage responses to 1 µM adenosine were observed in control, this agonist failed to hyperpolarize MSCs with both TRAM-34 and IbTX in the bath, although these blockers could not prevent Ca²⁺ mobilization (n = 3) (Fig. 7c). When applied alone, 1 μ M TRAM-34 suppressed voltage responses by 40–60% only (n = 3)(Fig. 7d, e). It thus appears that K_{Ca}3.1 and K_{Ca}1.1 channels are functionally co-expressed in ATP- and adenosineresponsive MSCs to couple agonist-dependent Ca2+ mobilization to cell hyperpolarization.

Discussion

Several lines of evidence point out that a branched Ca²⁺ signaling system involving multiple signaling pathways operates in MSCs and endows them with high capability to respond to environmental cues [19, 67]. In particular, Kawano and coworkers [31, 32] pioneered in demonstrating spontaneous Ca²⁺ oscillations in MSCs. The oscillatory behavior of intracellular Ca²⁺ in MSCs was associated with paracrine/ autocrine release of ATP that stimulated P2Y receptors coupled to phospholipase C (PLC) activation, IP₃ production, and liberation of Ca²⁺ ions from Ca²⁺ store. When enhanced by cyclic ADP-ribose via activation of TRPM2 channels, Ca²⁺ oscillation can stimulate MSC proliferation associated with enhanced ERK1/2 phosphorylation [60]. Reportedly, certain agonists of G-protein-coupled receptors, including adenosine, acetylcholine, ATP, histamine, noradrenaline, and some others, are capable of mobilizing intracellular Ca²⁺ in MSCs [18, 32, 35, 58]. Acetylcholine stimulates MSC migration by involving M1 receptors coupled to the PLC-PKC-ERK1/2 signaling pathway [58]. Functional and molecular evidence indicates that IP₃ receptors serve as dominant Ca²⁺ release channels in MSCs, while a role for ryanodine receptors in Ca²⁺ signaling in these cells remains obscure [32, 35]. In MSCs derived from the human adipose tissue, IP₃ receptors mediate Ca2+-induced Ca2+ release, a key intracellular event in adrenergic transduction [35]. Bath Ca²⁺ can influence the proliferation and osteogenic differentiation of MSCs from human bone marrow by stimulating the extracellular Ca²⁺-sensing receptor coupled to protein kinase C, thus initiating Ras-MAP kinase signaling [2]. Although voltage-gated (VG) Ca²⁺ channels are an attribute of excitable cells, several reports suggest that MSCs also employ these channels [1, 2, 40]. As is the case with apparently all excitable and non-excitable cells, store-operated channels are functional in MSCs as well [32]. The relaxation of Ca²⁺ transients in the MSC cytoplasm involves reticular and plasmalemmal Ca²⁺ ATPases and Na⁺/ Ca²⁺ exchange [31, 32].

By coupling intracellular Ca^{2+} signals to a change in transmembrane voltage, Ca^{2+} -dependent ion channels may contribute to cellular functions involving transport, receptor, and signaling proteins operating in the plasma membrane. The diverse family of Ca^{2+} -gated channels includes K⁺ channels of small ($K_{Ca}2.x$, formerly SK), intermediate ($K_{Ca}3.1$, IK), and large ($K_{Ca}1.1$, BK) conductance [4, 28, 38, 56]; cation channels TRPM4 and TRPM5 [21]; and anion channels [37, 53]. To our knowledge, the activity or expression of any Ca²⁺gated anion channel in MSCs has not been demonstrated yet, while functional expression of TRPM4 has been stated in a single report, wherein MSCs from the human adipose tissue were studied [62]. In contrast, multiple reports indicate that Ca²⁺-gated K⁺ channels operate in MSCs from various tissues of different species. In particular, evidence exists that



Fig. 6 Voltage and Ca²⁺ responses of MSCs to ATP. a Representative concurrent monitoring of intracellular Ca²⁺ (upper panel) and zerocurrent voltage (bottom panel) in MSC loaded with Fluo-4 (n = 9). Upper panel, the serial stimulation of a cell with 10 µM ATP elicited large Ca²⁺ transients presented as $\Delta F/F_0$. Bottom panel, voltage responses of the same cell to ATP were partly inhibited by 1 µM TRAM-34 but completely suppressed in the presence of 1 μ M TRAM-34 and 100 nM IbTX, suggesting those to be mediated by K_{Ca}1.1 and K_{Ca}3.1 channels. At the moments 1-12, I-V curves were generated by voltage ramps (1 mV/ms) from -100 to 80 mV. b Being generated in control (curve 1), during ATP application (curve 2) and after the rinse of ATP and Ca²⁺ relaxation (curve 3) (moments 1-3 in (a), respectively), the I-V curves 1-3 pointed out that an ATP-induced rise in intracellular Ca²⁺ was associated with a marked increase in inwardly rectifying conductance (left panel). Right panel, calculated as a difference between integral currents recorded in the presence of ATP and in control, an ATPstimulated current reversed below -80 mV (arrow). c Membrane voltages recorded in nine different cells in control, in the presence of ATP (the first application), and after rinse of the nucleotide. d I-V curves generated at the moments 4-12 indicated that outwardly

 K_{Ca} 1.1 channels are functionally expressed in bone marrowderived MSCs (BD-MSCs) from the human [24, 32], rabbit [16], and rat [24, 40, 64]. In addition to dominant K_{Ca} 1.1 channels, rat and rabbit BM-MSCs also express K_{Ca} 3.1

rectifying currents stimulated by ATP were completely and reversibly inhibited by 100 nM IbTX, pointing at K_{Ca}1.1 channels to be largely responsible. e Summary of Ca²⁺ and voltage responses of MSCs to 10 µM ATP under different conditions. To compare different experiments, each ATP response obtained in the presence of the particular blocker was normalized to a corresponding responses recorded in control. Here, $\Delta F = F_{ATP} - F_0$, where F_0 and F_{ATP} are intensities of cell fluorescence recorded just before ATP application and when intracellular Ca²⁺ peaked, $\Delta F_{\rm C}$ and $\Delta F_{\rm B}$ are Ca²⁺ responses recorded in control and in the presence of the particular blocker; $\Delta V = V_{ATP} - V_0$ where V_0 and V_{ATP} are the membrane voltages recorded before ATP application and when intracellular Ca²⁺ peaked, $\Delta V_{\rm C}$ and $\Delta V_{\rm B}$ are voltage responses recorded in control and in the presence of the particular blocker. The data are presented as a mean \pm SD (n = 9). The recording conditions were as in Fig. 1. In line with Student's unpaired ttest performed using SigmaPlot (Jandel Scientific), the difference between the control voltage response to ATP and responses recorded in the presence of TRAM-34 and TRAM-34 + IbTX is statistically significant (P < 0.05)

channels [16, 40, 64]. Mouse BD-MSCs appear to utilize solely K_{Ca} 3.1 channels. Indeed, among five genes coding for Ca^{2+} -gated K⁺ channels, only KCNN4 transcripts were identified in these cells. Yet, no IbTX-blockable currents were



Fig. 7 Voltage and Ca²⁺ responses of MSCs to adenosine. **a** Concurrent monitoring of intracellular Ca²⁺ (*upper panel*) and zero-current voltage (*bottom panel*) in MSC stimulated with 1 μ M adenosine. The shown traces were obtained by averaging recordings from eight cells and presented as a mean + SEM **b** Representative I-V curves that were generated in control during adenosine application and after the rinse of the agonist. *Insert*, adenosine-stimualated current, which is calculated as a difference between integral currents recorded in the presence of adenosine and in control, reverses below -80 mV (arrow). **c** Representative concurrent recording of intracellular Ca²⁺ (*upper panel*) and zero-current voltage (*bottom panel*) in MSC stimulated with 1 μ M adenosine in control, in the presence of 1 μ M TRAM-34 + 100 nM IbTX,

detected in mouse BD-MSCs but solely Ca^{2+} -activated K⁺ currents sensitive to clotrimazole, an K_{Ca}3.1 channel blocker [59]. K_{Ca}1.1 channels also were identified in MSCs from the human umbilical cord vein [43].

Here, we studied MSCs derived from the human adipose tissue (hAD-MSCs). The concurrent monitoring of cytosolic Ca^{2+} and membrane voltage from spontaneously oscillating hAD-MSCs provided the first evidence that intracellular Ca^{2+} signaling in these cells could be coupled to a change in membrane voltage (Fig. 1). Although in this particular case, spontaneous Ca^{2+} bursts elicited cell hyperpolarization by

and after rinse of the blockers (n = 3). **d** Representative concurrent recording of intracellular Ca²⁺ (*upper panel*) and zero-current voltage (*bottom panel*) in MSC stimulated with 1 μ M adenosine in control and in the presence of 1 μ M TRAM-34. **e** Summary of Ca²⁺ and voltage responses of MSCs to 1 μ M adenosine. Here, ΔF_C , ΔF_B , ΔV_C , and ΔV_B have the same meaning as in Fig. 6e, except for that adenosine responses instead of ATP responses are considered. The data are presented as a mean \pm SD (n = 3). The recording conditions were as in Fig. 1. The difference between the control voltage response to ATP and the responses recorded in the presence of TRAM-34 and TRAM-34 + IbTX is statistically significant (P < 0.05)

stimulating Ca²⁺-dependent K⁺ channels (Fig. 1), the question remained whether this coupling was universal for all MSCs or specific for the oscillating subpopulation. The expression analysis revealed mRNA transcripts for Ca²⁺-gated K⁺ channels of the K_{Ca}1.1 and K_{Ca}3.1 types as well as for some other Ca²⁺-activated channels, including the cationic channels TRPM4 and anionic channels from the anoctamin and bestrophin families (Fig. 2). Theoretically, this set of Ca²⁺gated ion channels could allow hAD-MSCs to generate diverse electrophysiological responses, depending on a combination of channel subunits expressed in a given cell and/or magnitude of Ca²⁺ signals triggered by external stimuli. When we treated hAD-MSCs with the Ca²⁺ ionophore ionomycin, all assayed cells were hyperpolarized by 20–40 mV (Fig. 3). This effect of artificially elevated cytosolic Ca²⁺ was mostly mediated by K_{Ca}1.1 channels, which appear to be ubiquitous for hAD-MSCs, although in a small (~12%) cellular subpopulation, K_{Ca}3.1 channels also contributed to voltage responses to ionomycin (Fig. 3d, e).

To our knowledge, electrophysiological properties of hAD-MSC were analyzed previously in a single work [1]. Based on functional and molecular data, the authors declared that AD-MSC expresses several families of ion channels, including VG Na⁺ channels and Ca²⁺-activated K⁺ channels of all three types. In particular, they detected transcripts for KCNMA1, KCNN3, and KCNN4 genes and recorded apamin (K_{Ca}2.x blocker)-, clotrimazole-, and IbTX-sensitive Ca²⁺-activated K⁺ currents in 83, 79, and 31% cellular fractions, respectively. These findings differ from our data in that we detected no Ca^{2+} -activated K⁺ currents sensitive to apamin (not shown). although more than 200 AD-MSCs were assayed overall. Consistently, we failed to detect transcripts for the KCNN1-KCNN3 genes encoding K_{Ca}2.x channels (Fig. 2a). Although at the moment, we cannot reasonably address this inconsistency; somewhat, different protocols of cell isolation and maintenance might result in rather different cellular compositions of AD-MSC populations assayed by us and in the work [1].

When assayed with perforated patch approach and CsCl in the patch pipette, every third cell responded to ionomycin with a marked inward current. This Ca2+-dependent current was negligibly sensitive to the specific TRPM4 blocker 9phenanthrol (Supplementary Materials, Fig. S1). Thus, despite the detection of TRPM4 transcripts in RNA isolated from hSD-MSC samples (Fig. 2), we failed to get convincing functional evidence for activity of TRPM4 channels. Perhaps, these channels were insufficiently active or not expressed in the vast majority of hSD-MSCs. The introduction of 440 nM Ca²⁺ into hSD-MSCs via the recording pipette stimulated anion currents in some of them (27%), and in all such cells, Ca²⁺-dependent currents were inhibited by CaCCinh completely. Because subtype-specific antagonists are not available, except for the Ano1 inhibitor T16, it was impossible to judge conclusively on the molecular identity of anionic channels responsible for Ca²⁺-gated currents. Note, however, that among hAD-MSCs responsive to Ca²⁺ loading, outwardly rectifying slowly activating currents appeared in every third cell as cytosolic Ca^{2+} rose during cell dialysis (Fig. 4). By activation kinetics and voltage dependence (Fig. 4), these currents resembled Ca²⁺-gated anion currents mediated by Ano1/ Ano2 channels [12, 46]. Since T16 exerted negligible effects, Ano2 channels were presumably responsible for these slow anion currents gated by cytosolic Ca²⁺ in hSD-MSCs. In nearly 70% fraction of Ca²⁺-sensitive cells, elevated cytosolic Ca²⁺ stimulated anion currents that were weakly rectifying and activated by electrical pulses almost instantly (Fig. 5), implicating bestrophin-like channels [41].

In a number of experiments, we analyzed whether Ca²⁺gated ion channels identified in hAD-MSCs might couple natural, that is elicited by agonists, Ca²⁺ signals to membrane polarization. The purinergic ligands ATP and adenosine were tested, because previously, we found both to mobilize Ca^{2+} in cells of this type [35]. It turned out that cells responsive, in terms of Ca^{2+} signaling, to 10 μ M ATP and 1 μ M adenosine were strongly hyperpolarized by both compounds (Figs. 6 and 7). The agonist-dependent hyperpolarization was partly diminished by TRAM-34 (1 µM) and completely disappeared when both TRAM-34 and IbTX (100 nM) were added to the bath (Figs. 6a and 7c). It thus appears that in hAD-MSCs responsive to ATP and adenosine, transduction of these agonists is associated with the generation of Ca²⁺ transients and activation of K⁺ channels of the K_{Ca}1.1 and K_{Ca}3.1 types. It is noteworthy that in some cells, hyperpolarizing voltage responses to the agonists were followed by conspicuous offresponses (Figs. 6a and 7d). This depolarizing phase of voltage responses might be mediated by Ca²⁺-gated anion channels, deactivation of which during relaxation of Ca^{2+} bursts was presumably delayed due to higher sensitivity to cytosolic Ca^{2+} compared to Ca^{2+} -dependent K⁺ channels.

In electrically excitable cells, K_{Ca}1.1 channels are usually co-localized with VG Ca²⁺ channels and control both the firing frequency and the action potential repolarization phase, thereby regulating neurotransmission in neurons, contraction in muscles, release of catecholamines in chromaffin cells, and insulin secretion in β cells [38, 63]. The physiological role of K_{Ca}1.1 and other Ca²⁺-gated channels identified in nonexcitable cells, including hAD-MSCs, remains poorly understood with few exceptions such as exocrine gland cells. In these cells, the acute regulation of fluid and electrolyte secretion depends on a fine control of K_{Ca}1.1 and VG K⁺ channels in conjunction with Ca²⁺-activated Cl⁻ channels [44–47]. Consistently with the expression of multiple P2Y and P1 purinoreceptors in MSCs of different origin, extracellular ATP and adenosine can modulate certain physiological processes in these cells [52]. Specifically, ATP can serve as adipogenic factor, while adenosine switches off adipogenic differentiation and promotes osteogenesis [10]. Our findings do not provide a clear idea of which cellular functions might depend on plasma membrane hyperpolarization elicited by ATP or adenosine and mediated by K_{Ca}1.1 and K_{Ca}3.1 channels in in hAD-MSCs. Note that proliferation and migration of diverse cells could be inhibited when activity of K_{Ca} 1.1 or K_{Ca}3.1 channels was suppressed by pharmacological blockage or genetic ablation [8, 20, 27, 36], although the underlying mechanisms were not clarified in most cases. The interesting possibility was proposed by Chandy and co-workers [9], who suggested that sequential activation of VG and Ca²⁺-activated K⁺ channels during mitosis triggers and sustains cell

hyperpolarization in T lymphocytes to facilitate Ca²⁺ entry via CRAC channels, thus stimulating the cell cycle machinery [9]. The somewhat similar mechanism was considered to explain the inhibitory effects of blockers of Ca2+-gated K+ channels on NO production stimulated by the Ca²⁺ mobilizing agents ATP and histamine in endothelial cells. It was suggested that by regulating Ca²⁺ influx responsible for stimulation of endothelial NO synthase, K_{Ca}2.x and K_{Ca}3.1 channel-mediated hyperpolarization represents a critical early event in agonist-evoked NO production [54]. Interestingly, NO has been reported as a MSC-derived modulator of T-cell proliferation in the mouse at least [51]. The similar hyperpolarization-dependent regulation of Ca²⁺ entry may take place in hAD-MSCs as a part of intracellular signaling stimulated by ATP and adenosine. Human MSCs secrete important regulatory molecules such as cytokines, growth factors, and hormones as well as microvesicles and exosomes containing not only peptides but also microRNA [55]. The non-constitutive Ca²⁺-dependent exocytosis in MSCs might vary with activity of K_{Ca}1.1 and K_{Ca}3.1 channels, which should modulate Ca2+ influx. Hopefully, further experiments will allow for verifying the physiological role of K_{Ca}1.1 and K_{Ca}3.1 channels in hAD-MSC functioning.

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